## **WEST Search History**

Hide Items	Restore	Clear	Cancel	CM
		1 - 1 - 1 - 1 - 1		3-20-07

DATE: Tuesday, March 20, 2007

Hide? <u>Set Name Query</u> $DB = PGPB, USPT, EPAB, JPAB, DWPI, PLUR = YES; OP = ADJ$			Hit Count
	L3	L2 same (mutation or polymorphism or variant or snp)	43.
	L2	(olfactory adj receptor) near5 (gene or nucleic or DNA or RNA or mRNA)	226
	L1	OR11H7p	1

END OF SEARCH HISTORY

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20mar07 14:32:52 User208746 Session D1147.2
                    0.115 DialUnits File410
            $0.00
     $0.00 Estimated cost File410
     $0.21 TELNET
     $0.21 Estimated cost this search
     $0.68 Estimated total session cost
                                          0.249 DialUnits
SYSTEM:OS - DIALOG OneSearch
  File
         5:Biosis Previews (R) 1926-2007/Mar W2
         (c) 2007 The Thomson Corporation
        5: BIOSIS has been enhanced with archival data. Please see
HELP NEWS 5 for information.
  File
         6:NTIS 1964-2007/Mar W3
         (c) 2007 NTIS, Intl Cpyrght All Rights Res
  File
         8:Ei Compendex(R) 1884-2007/Mar W1
         (c) 2007 Elsevier Eng. Info. Inc.
  File
        24:CSA Life Sciences Abstracts 1966-2007/Nov
         (c) 2007 CSA.
  File
        34:SciSearch(R) Cited Ref Sci 1990-2007/Mar W2
         (c) 2007 The Thomson Corp
  File
        45:EMCare 2007/Mar W2
         (c) 2007 Elsevier B.V.
  File
        65: Inside Conferences 1993-2007/Mar 20.
         (c) 2007 BLDSC all rts. reserv.
  File
        71:ELSEVIER BIOBASE 1994-2007/Mar W3
         (c) 2007 Elsevier B.V.
  File
        73:EMBASE 1974-2007/Mar 20
         (c) 2007 Elsevier B.V.
        94:JICST-EPlus 1985-2007/Mar W4
  File
         (c)2007 Japan Science and Tech Corp(JST)
*File 94: JICST will be removed from all vendors on March 31, 2007.
Please contact the Knowledge Center for alternative files.
  File 98:General Sci Abs 1984-2007/Mar
         (c) 2007 The HW Wilson Co.
        99:Wilson Appl. Sci & Tech Abs 1983-2007/Feb
         (c) 2007 The HW Wilson Co.
  File 135: NewsRx Weekly Reports 1995-2007/Mar W2
         (c) 2007 NewsRx
  File 136:BioEngineering Abstracts 1966-2007/Nov
         (c) 2007 CSA.
  File 143:Biol. & Agric. Index 1983-2007/Feb
         (c) 2007 The HW Wilson Co
  File 144:Pascal 1973-2007/Mar W2
         (c) 2007 INIST/CNRS
  File 155:MEDLINE(R) 1950-2007/Mar 15
         (c) format only 2007 Dialog
  File 172:EMBASE Alert 2007/Mar 20
         (c) 2007 Elsevier B.V.
  File 266:FEDRIP 2007/Feb
         Comp & dist by NTIS, Intl Copyright All Rights Res
  File 315: ChemEng & Biotec Abs 1970-2007/Feb
         (c) 2007 DECHEMA
  File 357:Derwent Biotech Res. 1982-2007/Mar W3
         (c) 2007 The Thomson Corp.
  File 358:Current BioTech Abs 1983-2006/Jan
         (c) 2006 DECHEMA
  File 369:New Scientist 1994-2007/Nov W3
         (c) 2007 Reed Business Information Ltd.
  File 370:Science 1996-1999/Jul W3
         (c) 1999 AAAS
*File 370: This file is closed (no updates). Use File 47 for more current
information.
  File 399:CA SEARCH(R) 1967-2007/UD=14613
         (c) 2007 American Chemical Society
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*File 399: Use is subject to the terms of your user/customer agreement.
IPCR/8 classification codes now searchable as IC=. See HELP NEWSIPCR.
 File 434:SciSearch(R) Cited Ref Sci 1974-1989/Dec
        (c) 2006 The Thomson Corp
     Set Items Description
? s OR11H7p
     S1 0 OR11H7P
? s olfactory adj receptor
     S2
          O OLFACTORY ADJ RECEPTOR
? s olfactory (w) receptor
         179905 OLFACTORY
        4805386 RECEPTOR
         17264 OLFACTORY (W) RECEPTOR
? s s3 and (polymorphism or snp or variant or mutation)
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         821294 POLYMORPHISM
          64258 SNP
         493944 VARIANT
        1636277 MUTATION
           616 S3 AND (POLYMORPHISM OR SNP OR VARIANT OR MUTATION)
? s s4 and py<2004
Processing
Processing
Processing
Processed 10 of 26 files ...
Processing
Processing
>>>One or more prefixes are unsupported
>>> or undefined in one or more files.
Processed 20 of 26 files ...
Processing
Completed processing all files
            616 S4
       127871174 PY<2004
     S5 463 S4 AND PY<2004
? rd s5
     S6
          293 RD S5 (unique items)
?
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     AAH31982;
XX
DT
     30-JUL-2001 (first entry)
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DE
     Human olfactory receptor polynucleotide, SEQ ID NO: 555.
XX
KW
     Human; olfactory receptor; OR; primary scent determination;
KW
     secondary scent determination; polypeptide library; odour receptor;
KW
     scent profile; scent fingerprint; scent representation; ds.
XX
OS
     Homo sapiens.
XX
PN
     WO200127158-A2.
XX
PD
     19-APR-2001.
XX
PF
     06-OCT-2000; 2000WO-US027582.
XX
PR
     08-OCT-1999;
                   99US-0158615P.
PR
     24-FEB-2000; 2000US-0184809P.
XX
PA
     (DIGI-) DIGISCENTS.
PA
     (YEDA ) YEDA RES & DEV CO LTD.
XX
PΙ
                  Smith D, Lancet D,
                                      Glusman G,
                                                  Fuchs T,
XX
DR
     WPI; 2001-290713/30.
XX
PT
     New polynucleotides which encode polypeptides involved in olfactory
PT
     sensation for identifying olfactory agonists and antagonists.
XX
PS
     Claim 8; Page 402; 1857pp; English.
XX
CC
     The present sequence is one of a number of isolated polynucleotides which
CC
     encode polypeptides involved in olfactory sensation. The polynucleotides
CC
     can be used in screening for olfactory agonists and antagonists. The
CC
     methods allow for the determination of primary scents and the
CC
     identification of the odour receptors used to detect these primary
CC
     scents. The methods also enable determination of secondary scents and the
CC
     identification of combinations of odour receptors that are involved in
CC
     detecting such secondary scents. This enables the construction of a scent
CC
     representation (also called a scent fingerprint or scent profile), which
CC
     may be used to re-create and edit scents. Libraries of olfactory
     receptors are useful for determining the interaction pattern of a
CC
CC
     composition with the receptors, and can be used for determining
CC
     differences in the olfactory faculties of different individuals
XX
SQ
     Sequence 960 BP; 203 A; 252 C; 184 G; 321 T; 0 U; 0 Other;
  Query Match
                         100.0%; Score 954; DB 4; Length 960;
  Best Local Similarity
                         100.0%; Pred. No. 5.2e-279;
 Matches 954; Conservative
                               0; Mismatches
                                                 0;
                                                     Indels
Qу
           1 AGTCTGGGAAGCATGAATAACTCACAGATATCTACTGTGACGCAGTTTGTGTTGTGGGG 60
             Db
           1 AGTCTGGGAAGCATGAATAACTCACAGATATCTACTGTGACGCAGTTTGTGTTGTTGGGG 60
Qу
          61 TTTCCTGGTCCCTGGAAAATTCAGATCATCTTTTTCTCAATGATTTTGTTGGTCTACATC 120
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Qy	181	ACCCCTATGTACGTGCTCCTAGCCAACTTCTCCTTCCTAGAGATCTGGTATGTGACCTGC	240
Db	181	ACCCCTATGTACGTGCTCCTAGCCAACTTCTCCTTCCTAGAGATCTGGTATGTGACCTGC	240
Qу	241	ACAGTCCCCAACATGCTGGTAAATTTTTTCTCCAAAACTAAGACCATATCATTCTCTGGA	300
Db	241	ACAGTCCCCAACATGCTGGTAAATTTTTTCTCCAAAACTAAGACCATATCATTCTCTGGA	300
Qγ	301	TGTTTCACTCAGTTCCACTTCTTCTTTTCCCTGGGCACAACTGAATGCTTCTTCCTCTGT	360
Db	301	TGTTTCACTCAGTTCCACTTCTTTTTCCCTGGGCACAACTGAATGCTTCTTCCTCTGT	360
Qy	361	GTCATGGCTTATGATCGGTACCTGGCCATCTGCCACCCAC	420
Db	361	GTCATGGCTATGATCGGTACCTGGCCATCTGCCACCCACTGCACTATCCCTCCATTATG	420
Qy	421	ACTGGCCAGCTCTGTGGCATCTTGGTGTCTCTTTGTTGGCTCATTGGTTTCCTTGGACAT	480
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Qy	481	TCAATTTCCATTTTCTTCATTTTTCAACTACCTTTCTGTGGTCCCAACATCATTGATCAT	540
Dр	481	TCAATTTCCATTTTTCAACTACCTTTCTGTGGTCCCAACATCATTGATCAT	540
Qy	541	TTTCTGTGTGATGTAGACCCACTGATGGCATTGTCCTCTGCCCCTACTCACATCATAGGG	600
Db	541	TTTCTGTGTGATGTAGACCCACTGATGGCATTGTCCTCTGCCCCTACTCACATCATAGGG	600
Qy		CATGTGTTCCATTCTGTGAGCTCTCTTTTCATCAACCTCACCATGGTGTACATCCTTGGG	660
Db		CATGTGTTCCATTCTGTGAGCTCTCTTTTCATCAACCTCACCATGGTGTACATCCTTGGG	660
Qу	661	TCCTATACCTTGGTGCTCAGAACTGTGCTTTAGGTTCCTTCTTCAGCTGGATGGCAAAAG	720
Db	661	TCCTATACCTTGGTGCTCAGAACTGTGCTTTAGGTTCCTTCTTCAGCTGGATGGCAAAAG	720
Qy	721	GCCATCTCTACCTGTGGGTCACACTTGGTTGTTGTGTCTCTGTTCTATGGAGCCATAATG	780
Db	721	GCCATCTCTACCTGTGGGTCACACTTGGTTGTTGTTGTTCTTATGGAGCCATAATG	780
Qy	781	CTGATGTATGTGAGTCCCACACCTGGCAACTCAGTTGCTATGCATAAGCTCATCACACTG	84,0
Db	781	CTGATGTATGTGAGTCCCACACCTGGCAACTCAGTTGCTATGCATAAGCTCATCACACTG	840
Qy	841	ATATATTCTGTGGTAACACCTGTCTTAAACCCCCTCATCTACAGCCTACGCAACAAGGAC	900
Db	841	ATATATTCTGTGGTAACACCTGTCTTAAACCCCCTCATCTACAGCCTACGCAACAAGGAC	900
Qy	901	ATGAAATATGCCCTCCATCATGTCTTCTGTGGAATGAGAATTATCCAGAGATCA 954	
Db EndFra</td <td></td> <td>ATGAAATATGCCCTCCATCATGTCTTCTGTGGAATGAGAATTATCCAGAGATCA 954</td> <td></td>		ATGAAATATGCCCTCCATCATGTCTTCTGTGGAATGAGAATTATCCAGAGATCA 954	